

Lactarius deliciosus isolate UEZB1 internal transcribed spacer 1, partial sequence; 5.8S ribosomal RNA gene and internal transcribed spacer 2, complete sequence; and 28S ribosomal RNA gene, partial sequence

GenBank: JQ066791.1

[FASTA Graphics](#)

[Go to:](#)

LOCUS JQ066791 439 bp DNA linear PLN
20-MAR-2012
DEFINITION Lactarius deliciosus isolate UEZB1 internal transcribed
spacer 1,
partial sequence; 5.8S ribosomal RNA gene and internal
transcribed
spacer 2, complete sequence; and 28S ribosomal RNA gene,
partial
sequence.
ACCESSION JQ066791
VERSION JQ066791.1 GI:380467948
KEYWORDS .
SOURCE Lactarius deliciosus
ORGANISM [Lactarius deliciosus](#)
Eukaryota; Fungi; Dikarya; Basidiomycota; Agaricomycotina;
Agaricomycetes; Russulales; Russulaceae; Lactarius.
REFERENCE 1 (bases 1 to 439)
AUTHORS Ragonezi,C., Caldeira,A.T., Martins,M.R., Santos-Silva,C.,
Klimaszewska,K., Louro,R., Ganhao,E. and Zavattieri,A.
TITLE Direct Submission
JOURNAL Submitted (23-NOV-2011) Chemistry Department, University
of Evora,
Rua Romao Ramalho, 59, Evora 7000 671, Portugal
FEATURES Location/Qualifiers
source 1..439
/organism="Lactarius deliciosus"
/mol_type="genomic DNA"
/isolate="UEZB1"
/db_xref="taxon:[55514](#)"
[misc_RNA](#) <1..14
/product="internal transcribed spacer 1"
[rRNA](#) 15..172
/product="5.8S ribosomal RNA"
[misc_RNA](#) 173..381
/product="internal transcribed spacer 2"
[rRNA](#) 382..>439
/product="28S ribosomal RNA"
ORIGIN

1 acgcgcaatc aatacaactt tcaacaacgg atctcttggc tctcgcatcg
atgaagaacg
61 cagcgaaatg cgatacgtaa tgtgaattgc agaattcagt gaatcatcga
atctttgaac
121 gcaccttgcg ccccttggtg ttccgagggg cacaccegtt tgagtgtcgt
gaaattctca
181 accttctcgg ttttcttctg gacaccgaag gaggcttgga cattggagggc
ctttgctggc
241 gtctcttaga cccagctcct cttaaataaa ttagcggggg cctctttgcc
gatccttgac
301 atgtgataag atgtttccat gacttgggtt ctggctctgt tgcatttggg
acccgcttct
361 aaccgtctcg acgagacaac gtttgggcgt gtctcccttc tcgggagact
ctctcaaccc
421 ccacgaaccc ttgacctca